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Reviewer: Durreshwar Anjum

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Application No: 10528104 Version No: 2.0

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SEQUENCE LISTING

<110> HEINZEL, THORSTEN
 KRAEMER, OLIVER H.
 GOETTLICHER, MARTIN
 ZHU, PING
 GOLEBIEWSKI, MARTIN
 PELICCI, PIER
 MAURER, ALEXANDER
 HENTSCH, BERND
 MINUCCI, SAVERIO

<120> THE USE OF MOLECULAR MARKERS FOR THE PRECLINICAL AND
 CLINICAL PROFILING OF INHIBITORS OF ENZYMES HAVING
 HISTONE DEACETYLASE ACTIVITY

<130> LEDER-15

<140> 10528104

<141> 2005-09-28

<150> PCT/EP03/10404

<151> 2003-09-18

<150> EP 02021228.8

<151> 2002-09-18

<160> 16

<170> PatentIn Ver. 3.3

<210> 1

<211> 488

<212> PRT

<213> Homo sapiens

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 35 40 45

Tyr Arg Lys Met Glu Ile Tyr Arg Pro His Lys Ala Thr Ala Glu Glu
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Met Thr Lys Tyr His Ser Asp Glu Tyr Ile Lys Phe Leu Arg Ser Ile
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Arg Pro Asp Asn Met Ser Glu Tyr Ser Lys Gln Met His Ile Phe Asn
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Tyr Phe Gly Pro Asp Phe Lys Leu His Ile Ser Pro Ser Asn Met Thr				
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Asn Gln Asn Thr Pro Glu Tyr Met Glu Lys Ile Lys Gln Arg Leu Phe				
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<212> PRT

<213> Homo sapiens

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35 40 45

Asp Asn Asn Leu Leu Gly Thr Pro Gly Glu Ser Thr Glu Glu Glu Leu
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Leu Arg Arg Leu Gln Gln Ile Lys Glu Gly Pro Pro Pro Gln Asn Ser
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Ser Ile Ile Asp Trp Leu Asn Ser Val Arg Gln Thr Gly Asn Thr Thr
100 105 110

Arg Ser Gly Gln Arg Gly Asn Gln Ser Trp Arg Ala Val Cys Arg Thr
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Asn Pro Asn Ser Gly Asn Phe Arg Phe Ser Leu Glu Ile Asn Val Tyr
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Ser Asn Asn Gly Ser Gln Asn Ser Glu Asn Glu Asn Glu Pro Ser Ala
145 150 155 160

Arg Arg Ser Ser Gly Glu Asn Val Glu Asn Asn Ser Gln Arg Gln Val
165 170 175

Glu Asn Pro Arg Ser Glu Ser Thr Ser Ala Arg Pro Ser Arg Ser Glu
180 185 190

Arg Asn Ser Thr Glu Ala Leu Thr Glu Val Pro Pro Thr Arg Gly Gln
195 200 205

Arg Arg Ala Arg Ser Arg Ser Pro Asp His Arg Arg Thr Arg Ala Arg
210 215 220

Ala Glu Arg Ser Arg Ser Pro Leu His Pro Met Ser Glu Ile Pro Arg
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Arg Ser His His Ser Ile Ser Ser Gln Thr Phe Glu His Pro Leu Val
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Asp	Ser	Ile	Ala	Ser	Arg	Thr	Arg	Ser	Arg	Ser	Gln	Thr	Pro	Asn	Asn	
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Thr	Val	Thr	Tyr	Glu	Ser	Glu	Arg	Gly	Gly	Phe	Arg	Arg	Thr	Phe	Ser	
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Ser	Ser	Ser	Ser	Ser	Ser	Ser	Ser	Ser	Ser	Ser	Ser	Pro	Ser	Ser	Ser	
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Tyr Thr Glu Gly Asn Lys Leu Arg Lys Leu Pro Cys Ser His Glu Tyr
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Ile Cys Arg Arg Ala Val Leu Ala Ser Gly Asn Arg Glu Ser Val Val
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<211> 281

<212> PRT

<213> Homo sapiens

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Val Thr Tyr Val Tyr Phe Thr Asn Glu Leu Lys Gln Met Gln Asp Lys
35 40 45

Tyr Ser Lys Ser Gly Ile Ala Cys Phe Leu Lys Glu Asp Asp Ser Tyr
50 55 60

Trp Asp Pro Asn Asp Glu Glu Ser Met Asn Ser Pro Cys Trp Gln Val
65 70 75 80

Lys Trp Gln Leu Arg Gln Leu Val Arg Lys Met Ile Leu Arg Thr Ser
85 90 95

Glu Glu Thr Ile Ser Thr Val Gln Glu Lys Gln Gln Asn Ile Ser Pro
100 105 110

Leu Val Arg Glu Arg Gly Pro Gln Arg Val Ala Ala His Ile Thr Gly
115 120 125

Thr Arg Gly Arg Ser Asn Thr Leu Ser Ser Pro Asn Ser Lys Asn Glu
130 135 140

Lys Ala Leu Gly Arg Lys Ile Asn Ser Trp Glu Ser Ser Arg Ser Gly
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His Ser Phe Leu Ser Asn Leu His Leu Arg Asn Gly Glu Leu Val Ile
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His Glu Lys Gly Phe Tyr Tyr Ile Tyr Ser Gln Thr Tyr Phe Arg Phe

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185

190

Gln Glu Glu Ile Lys Glu Asn Thr Lys Asn Asp Lys Gln Met Val Gln
 195 200 205

Tyr Ile Tyr Lys Tyr Thr Ser Tyr Pro Asp Pro Ile Leu Leu Met Lys
 210 215 220

Ser Ala Arg Asn Ser Cys Trp Ser Lys Asp Ala Glu Tyr Gly Leu Tyr
 225 230 235 240

Ser Ile Tyr Gln Gly Gly Ile Phe Glu Leu Lys Glu Asn Asp Arg Ile
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<212> DNA

<213> Homo sapiens

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atgagggttac gatcctggga ggacttaatg aatttgtagt gaagttttat ggaccacaag 180
gaacaccata tgaaggcgga gtatggaaaag ttagagtgga cctacctgat aaataccctt 240
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caggaactgt gtgtctagat gtaattaatc aaacttggac agctctctat gatcttacca 360
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agtacatcca gaaatacgcc acggaggagg cgctgaaaga acaggaagag ggtaccgggg 540
acagctcatc ggagagctct atgtctgact tttccgaaga tgaggcccag gatatggagt 600
tgtagtagaa aaagcacctg cttttcagaa agactattat ttcctaacca tgagaagcag 660
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